SCORE Search Results Details for Application 10660811 and Search Result 20061017_105802_us-10-660-811a-141.100aligns.rapbm.

Score Home Retrieve Application SCORE System SCORE Comments / Suggestions Page List Overview FAQ

This page gives you Search Results detail for the Application 10660811 and Search Result 20061017_105802_us-10-660-811a-141,100aligns,rapbm.

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OM protein - protein search, using sw model

Run on:

October 17, 2006, 16:29:12; Search time 184 Seconds

(without alignments)

2728.940 Million cell updates/sec

Title:

US-10-660-811A-141

Perfect score: 5550

Sequence:

1 VKSASALEHDNKLLLKWTTL.....EIGEVLVHIEQALEFVRKTG 1084

Scoring table:

BLOSUM62

Gapop 10.0, Gapext 0.5

Searched:

2097797 segs, 463214858 residues

Total number of hits satisfying chosen parameters: - 2097797

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 300 summaries

Database: Published_Applications_AA_Main:*

/EMC_Celerra_SIDS3/ptodata/2/pubpaa/US07_PUBCOMB.pep:*
/EMC_Celerra_SIDS3/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
/EMC_Celerra_SIDS3/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
/EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*
/EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
/EMC_Celerra_SIDS3/ptodata/2/pubpaa/US11_PUBCOMB.pep:* 3: 4:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
. 1 2 3	5550 2685 2680	100.0 48.4 48.3	1084 664 663	5	US-10-660-811A-141 US-10-482-948-6 US-10-482-948-35	Sequence 141, App Sequence 6, Appli Sequence 35, Appl

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